http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?db=protein&qty=1&...uids=NP\_001142.2&dopt=fasta&dispmax=5&sendto=t&from=be; ANT-1 AS DRUG TARGET

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ANT-1 AS DRUG TARGET
Attorney Docket: 2923-478

>gi|55749577|ref|NP\_001142.2| solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 4 [Homo sapiens]
MGDHAWSFLKDFLAGGVAAAVSKTAVAPIERVKLLLQVQHASKQISAEKQYKGIIDCVVRIPKEQGFLSF
WRGNLANVIRYFPTQALNFAFKDKYKQLFLGGVDRHKQFWRYFAGNLASGGAAGATSLCFVYPLDFARTR
LAADVGKGAAQREFHGLGDCIIKIFKSDGLRGLYQGFNVSVQGIIIYRAAYFGVYDTAKGMLPDPKNVHI
FVSWMIAQSVTAVAGLVSYPFDTVRRRMMMQSGRKGADIMYTGTVDCWRKIAKDEGAKAFFKGAWSNVLR
GMGGAFVLVLYDEIKKYV

http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?db=nucleotide&qty=...ist\_uids=55749576&dopt=fasta&dispmax=5&sendto=t&from=begin&to= Application No. 10/069,973

Application No. 10/069,973 ANT-1 AS DRUG TARGET Attorney Docket: 2923-478

>gi|55749576|ref|NM\_001151.2| Homo sapiens solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 4 (SLC25A4), nuclear gene encoding mitochondrial protein, mRNA

GGGGAGCTGCGGGCCAGGCGCCCCCTAGCGTCGCGCAGGGTCGGGGACTGCGCGGCGGTGCCAGGC CGGGCGTGGGCGAGAGCACGAACGGGCTGCCGGGCTGAGAGCGTCGAGCTGTCACCATGGGTGATCA CGCTTGGAGCTTCCTAAAGGACTTCCTGGCCGGGGGGGCGTCGCCGCTGCCGTCTCCAAGACCGCGGTCGCC CCCATCGAGAGGGTCAAACTGCTGCTGCAGGTCCAGCATGCCAGCAAACAGATCAGTGCTGAGAAGCAGT ACAAAGGGATCATTGATTGTGTGGTGAGAATCCCTAAGGAGCAGGGCTTCCTCCTCTCTGGAGGGGTAA CCTGGCCAACGTGATCCGTTACTTCCCCACCCAAGCTCTCAACTTCGCCTTCAAGGACAAGTACAAGCAG CTCTTCTTAGGGGGTGTGGATCGGCATAAGCAGTTCTGGCGCTACTTTGCTGGTAACCTGGCGTCCGGTG GGGCCGCTGGGGCCACCTCCCTTTGCTTTGTCTACCCGCTGGACTTTGCTAGGACCAGGTTGGCTGCTGA TGTGGGCAAGGGCGCCCAGCGTGAGTTCCATGGTCTGGGCGACTGTATCATCAAGATCTTCAAGTCT GATGGCCTGAGGGGGCTCTACCAGGGTTTCAACGTCTCTGTCCAAGGCATCATTATCTATAGAGCTGCCT ACTTCGGAGTCTATGATACTGCCAAGGGGATGCTGCCTGACCCCAAGAACGTGCACATTTTTGTGAGCTG GATGATTGCCCAGAGTGTGACGGCAGTCGCAGGGCTGGTGTCCTACCCCTTTGACACTGTTCGTCGTAGA ATGATGATGCAGTCCGGCCGGAAAGGGGCCGATATTATGTACACGGGGACAGTTGACTGCTGGAGGAAGA TTGCAAAAGACGAAGGACCAAGGCCTTCTTCAAAGGTGCCTGGTCCAATGTGCTGAGAGGCATGGGCGG TGCTTTTGTATTGGTGTTGTATGATGAGATCAAAAAATATGTCTAATGTAATTAAAACACAAGTTCACAG ATTTACAGTGAACTTGATCTACAAGTTCACAGATCCATTGTGTGGTTTAATAGACTATTCCTAGGGGAAG TAAAAAGATCTGGGATAAAACCAGACTGAAAGGAATACCTCAGAAGAGATGCTTCATTGAGTGTTCATTA AACCACACATGTATTTTGTATTTTATTTTACATTTAAATTCCCACAGCAAATAGAAAATAATTTATCATAC TTGTACAATTAACTGAAGAATTGATAATAACTGAATGTGAAACATCAATAAAGACCACTTAATGCAAAAA AAAAAAAA